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110/02

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/023,681

DATE: 09/10/2002 TIME: 15:32:46

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3 <110> APPLICANT: Li, Li
              Ballinger, Robert A
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              Padigaru, Muralidhara
              Kekuda, Ramesh
              Colman, Steven D
              Spytek, Kimberly A
      9
              Casman, Stacie J
     10
              Vernet, Corine AM
     11
              Shenoy, Suresh G
              Gusev, Vladimir Y
     12
     13
              Malyankar, Uriel M
     14
              Edinger, Shlomit R
     15
              Gerlach, Valerie
     16
              Smithson, Glennda
              Stone, David J
     17
     18
              Sciore, Paul
     19
              MacDougall, John R
     20
              Gunther, Erik
     21
              Peyman, John A
     22
              Ellerman, Karen
              Gangolli, Esha A
     23
     24
              Millet, Isabelle
     26 <120> TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
     28 <130> FILE REFERENCE: 21402-224AC
     30 <140> CURRENT APPLICATION NUMBER: 10/023,681
C--> 31 <141> CURRENT FILING DATE: 2002-09-04
     33 <150> PRIOR APPLICATION NUMBER: 60/256,635
     34 <151> PRIOR FILING DATE: 2000-12-18
     36 <150> PRIOR APPLICATION NUMBER: 60/259,743
     37 <151> PRIOR FILING DATE: 2001-01-04
     39 <150> PRIOR APPLICATION NUMBER: 60/299,327
     40 <151> PRIOR FILING DATE: 2001-06-19
     42 <150> PRIOR APPLICATION NUMBER: 60/261,498
     43 <151> PRIOR FILING DATE: 2001-01-12
     45 <150> PRIOR APPLICATION NUMBER: 60/263,689
     46 <151> PRIOR FILING DATE: 2001-01-24
     48 <150> PRIOR APPLICATION NUMBER: 60/267,464
     49 <151> PRIOR FILING DATE: 2001-02-08
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     54 <150> PRIOR APPLICATION NUMBER: 60/275,946
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73 <151> PRIOR FILING DATE: 2001-04-18
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87 acctdctcat cttqctqqct attgtctcct ctqctcgqct tcacacqccc atqtacttct 180
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92 teatectage catggggace teattettig getgeattea ggecaectit etgaceacte 480
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115 Gly Ile Tyr Val Val Thr Val Val Gly Asn Leu Ser Met Ile Thr Leu
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Input Set : A:\Cur524al.app

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130 His Met Leu Ala Ala Met Ala Tyr Asp His Tyr Ile Ala Ile Cys Asn
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133 Pro Leu Leu Tyr His Val Ala Met Ser Tyr Gln Val Cys Ser Trp Met
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136 Val Val Glu Val Tyr Phe Met Gly Phe Ile Gly Ala Thr Cys Ser His
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137 145
139 Ser Leu His Ala Lys Ser Ala Phe Leu Glu Gly Arg Cys Asn Gln Pro
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142 Leu Leu Gly Ser Phe Pro Thr Thr Gly Ala Leu Pro Leu Gln Tyr
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145 Phe Tyr Gln Arg Asn Ser Ser Leu Cys Phe Ser Ala Phe Asn Ile Leu
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                                                     205
148 Phe Arg Ser Leu Thr Ile Leu Ser Ser Tyr Ile Phe Ile Val Ala Ser
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151 Ile Leu Cys Ile Arg Ser Thr Glu Gly Arg Ser Lys Thr Phe Ser Thr
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152 225
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154 Cys Ser Ser His Ile Ser Ala Val Ser Val Phe Phe Gly Ser Ala Ala
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157 Phe Met Tyr Leu Gln Pro Ser Ser Val Ser Ser Met Asp Gln Gly Ser
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160 Val Phe Cys Val Leu Cys Tyr Cys Cys Ala His Ala Glu Pro Pro Ile
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178 acctgctcat cttgctggct attgtctcct ctgctcggct tcacacgccc atgtacttct 180
179 teetgtgeaa getgtetgtt tittgaeetat titteeette tgtgagttee eetaagatge 240
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181 tettetacea ttteetggge tgeactgagt gttteetgta caeggtgatg geetaegaee 360
182 gctttgttgc catttgtcac cctctacgct acaccataat catgagccac agagcatgta 420
183 teatectage catggggace teattettig getgeattea ggecacetti etgaceaete 480
184 teacetteca attgeettae tgtgteecea atgaggtgga etattattte tgtgatatee 540
185 cagteatget gaagetgget tgtgeagata ceteageeet ggagatggtg gggtteatea 600
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187 tettetecat ettggagate tgetetgeeg agggeegaeg eegtgeette teeaeetgea 720
188 gegeecacct cacegocate etgetttttt acatgocagt ggteeteatt tacetgagge 780
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206 Tyr Ile Phe Thr Leu Met Gly Asn Leu Leu Ile Leu Leu Ala Ile Val
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                                                    45
209 Ser Ser Ala Arg Leu His Thr Pro Met Tyr Phe Phe Leu Cys Lys Leu
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                                                 60
212 Ser Val Phe Asp Leu Phe Phe Pro Ser Val Ser Ser Pro Lys Met Leu
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215 Cys Tyr Leu Ser Gly Asn Ser Arg Ala Ile Ser Tyr Ala Gly Cys Ala
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218 Ser Gln Leu Phe Phe Tyr His Phe Leu Gly Cys Thr Glu Cys Phe Leu
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221 Tyr Thr Val Met Ala Tyr Asp Arg Phe Val Ala Ile Cys His Pro Leu
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224 Arg Tyr Thr Ile Ile Met Ser His Arg Ala Cys Ile Ile Leu Ala Met
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227 Gly Thr Ser Phe Phe Gly Cys Ile Gln Ala Thr Phe Leu Thr Thr Leu
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230 Thr Phe Gln Leu Pro Tyr Cys Val Pro Asn Glu Val Asp Tyr Tyr Phe
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233 Cys Asp Ile Pro Val Met Leu Lys Leu Ala Cys Ala Asp Thr Ser Ala
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236 Leu Glu Met Val Gly Phe Ile Ser Val Gly Leu Met Pro Leu Ser Cys
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239 Phe Leu Leu Ile Leu Thr Ser Tyr Ser Gly Ile Val Phe Ser Ile Leu
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242 Glu Ile Cys Ser Ala Glu Gly Arg Arg Ala Phe Ser Thr Cys Ser
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245 Ala His Leu Thr Ala Ile Leu Leu Phe Tyr Met Pro Val Val Leu Ile
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248 Tyr Leu Arg Pro Thr His Ser Leu Trp Leu Asp Ala Thr Val Gln Ile
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251 Leu Asn Asn Leu Val Thr Pro Met Leu Asn Pro Leu Ile Tyr Ser Leu
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269 acctgctcat cttgctggct attgtctcct ctgctcggct tcacacgccc atgtacttct 180
270 teetgtgeaa getgtetgtt tttgacetat tttteeette tgtgagttee cetaagatge 240
271 tgtgctatct ttcagggaac agccgagcca tctcctatgc aggctgtgca tcccagctct 300
272 tettetacca ttteetggge tgeactgagt gttteetgta caeggtgatg geetacgace 360
273 getttgttgc catttgtcac cetetacget acaccataat catgagecac agageatgta 420
274 teatectage catggggace teattettig getgeattea ggecacetti etgaceaete 480
275 teacetteca attgeettae tgtgteecca atgaggtgga etattattte tgtgatatee 540
276 cagtcatgct gaagctggct tgtgcagata cctcagccct ggagatggtg gggttcatca 600
277 gtgtgggcct catgcccctc agctgtttcc ttctcatcct cacctcctac agtggcatcg 660
278 tettetecat ettggagate tgetetgeeg agggeegaeg eegtgeette teeacetgea 720
279 gegeceacet cacegocate etgetttttt acatgocagt ggteeteatt tacetgagge 780
280 ctacccacag cctgtggttg gatgcaactg ttcaaattct gaataacctg gtcaccccca 840
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294 His Thr Glu Gly Leu Glu Thr Ile Leu Leu Val Leu Phe Leu Ser Phe
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297 Tyr Ile Phe Thr Leu Met Gly Asn Leu Leu Ile Leu Leu Ala Ile Val
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300 Ser Ser Ala Arg Leu His Thr Pro Met Tyr Phe Phe Leu Cys Lys Leu
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303 Ser Val Phe Asp Leu Phe Phe Pro Ser Val Ser Fro Lys Met Leu
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306 Cys Tyr Leu Ser Gly Asn Ser Arg Ala Ile Ser Tyr Ala Gly Cys Ala
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309 Ser Gln Leu Phe Phe Tyr His Phe Leu Gly Cys Thr Glu Cys Phe Leu
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312 Tyr Thr Val Met Ala Tyr Asp Arg Phe Val Ala Ile Cys His Pro Leu
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315 Arg Tyr Thr Ile Ile Met Ser His Arg Ala Cys Ile Ile Leu Ala Met
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318 Gly Thr Ser Phe Phe Gly Cys Ile Gln Ala Thr Phe Leu Thr Thr Leu
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321 Thr Phe Gln Leu Pro Tyr Cys Val Pro Asn Glu Val Asp Tyr Tyr Phe
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324 Cys Asp Ile Pro Val Met Leu Lys Leu Ala Cys Ala Asp Thr Ser Ala
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VERIFICATION SUMMARY PATENT APPLICATION: US/10/023,681 TIME: 15:32:47

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